



OIPE

RAW SEQUENCE LISTING  
PATENT APPLICATION: US/10/044,674

DATE: 02/11/2002  
TIME: 10:00:57

Input Set : A:\TNFRSF11B seq list.txt  
Output Set: N:\CRF3\02112002\J044674.raw

3 <110> APPLICANT: Chew, Anne  
4 Denton, R. Rex  
5 Bieglecki, Karyn M  
6 Nandabalan, Krishnan  
7 Stephens, J. Claiborne  
9 <120> TITLE OF INVENTION: HAPLOTYPES OF THE TNFRSF11B GENE  
11 <130> FILE REFERENCE: TNFRSF11B\_MWH-0001US (CIP)  
C--> 13 <140> CURRENT APPLICATION NUMBER: US/10/044,674  
14 <141> CURRENT FILING DATE: 2002-01-09  
16 <150> PRIOR APPLICATION NUMBER: PCT/US00/18803  
17 <151> PRIOR FILING DATE: 2000-07-10  
19 <160> NUMBER OF SEQ ID NOS: 94  
21 <170> SOFTWARE: PatentIn version 3.1

Does Not Comply  
Corrected Diskette Needed

ERRORED SEQUENCES

1143 <210> SEQ ID NO: 94  
1144 <211> LENGTH: 2280  
1145 <212> TYPE: DNA  
1146 <213> ORGANISM: Homo sapiens  
1148 <220> FEATURE:  
1149 <221> NAME/KEY: allele  
1150 <222> LOCATION: (30)..(30)  
1151 <223> OTHER INFORMATION: PS1: polymorphic base G or T  
1154 <220> FEATURE:  
1155 <221> NAME/KEY: misc\_feature  
1156 <222> LOCATION: (61)..(120)  
1157 <223> OTHER INFORMATION: n's represent sequence between PS1 and PS2  
1160 <220> FEATURE:  
1161 <221> NAME/KEY: allele  
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1163 <223> OTHER INFORMATION: PS2: polymorphic base C or T  
1166 <220> FEATURE:  
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1168 <222> LOCATION: (181)..(240)  
1169 <223> OTHER INFORMATION: n's represent sequence between PS2 and PS3  
1172 <220> FEATURE:  
1173 <221> NAME/KEY: allele  
1174 <222> LOCATION: (270)..(270)  
1175 <223> OTHER INFORMATION: PS3: polymorphic base G or T  
1178 <220> FEATURE:  
1179 <221> NAME/KEY: misc\_feature

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1180 <222> LOCATION: (301)..(360)  
1181 <223> OTHER INFORMATION: n's represent sequence between PS3 and PS4  
1184 <220> FEATURE:  
1185 <221> NAME/KEY: allele  
1186 <222> LOCATION: (390)..(390)  
1187 <223> OTHER INFORMATION: PS4: polymorphic base T or C  
1190 <220> FEATURE:  
1191 <221> NAME/KEY: misc\_feature  
1192 <222> LOCATION: (421)..(480)  
1193 <223> OTHER INFORMATION: n's represent sequence between PS4 and PS5  
1196 <220> FEATURE:  
1197 <221> NAME/KEY: allele  
1198 <222> LOCATION: (510)..(510)  
1199 <223> OTHER INFORMATION: PS5: polymorphic base G or C  
1202 <220> FEATURE:  
1203 <221> NAME/KEY: misc\_feature  
1204 <222> LOCATION: (541)..(600)  
1205 <223> OTHER INFORMATION: n's represent sequence between PS5 and PS6  
1208 <220> FEATURE:  
1209 <221> NAME/KEY: allele  
1210 <222> LOCATION: (630)..(630)  
1211 <223> OTHER INFORMATION: PS6: polymorphic base C or T  
1214 <220> FEATURE:  
1215 <221> NAME/KEY: misc\_feature  
1216 <222> LOCATION: (661)..(720)  
1217 <223> OTHER INFORMATION: n's represent sequence between PS6 and PS7  
1220 <220> FEATURE:  
1221 <221> NAME/KEY: allele  
1222 <222> LOCATION: (750)..(750)  
1223 <223> OTHER INFORMATION: PS7: polymorphic base G or A  
1226 <220> FEATURE:  
1227 <221> NAME/KEY: misc\_feature  
1228 <222> LOCATION: (781)..(840)  
1229 <223> OTHER INFORMATION: n's represent sequence between PS7 and PS8  
1232 <220> FEATURE:  
1233 <221> NAME/KEY: allele  
1234 <222> LOCATION: (870)..(870)  
1235 <223> OTHER INFORMATION: PS8: polymorphic base C or A  
1238 <220> FEATURE:  
1239 <221> NAME/KEY: misc\_feature  
1240 <222> LOCATION: (901)..(960)  
1241 <223> OTHER INFORMATION: n's represent sequence between PS8 and PS9  
1244 <220> FEATURE:  
1245 <221> NAME/KEY: allele  
1246 <222> LOCATION: (990)..(990)  
1247 <223> OTHER INFORMATION: PS9: polymorphic base C or T  
1250 <220> FEATURE:  
1251 <221> NAME/KEY: misc\_feature  
1252 <222> LOCATION: (1021)..(1080)

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1253 <223> OTHER INFORMATION: n's represent sequence between PS9 and PS10  
1256 <220> FEATURE:  
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1258 <222> LOCATION: (1110)..(1110)  
1259 <223> OTHER INFORMATION: PS10: polymorphic base T or C  
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1263 <221> NAME/KEY: misc\_feature  
1264 <222> LOCATION: (1141)..(1200)  
1265 <223> OTHER INFORMATION: n's represent sequence between PS10 and PS11  
1268 <220> FEATURE:  
1269 <221> NAME/KEY: allele  
1270 <222> LOCATION: (1230)..(1230)  
1271 <223> OTHER INFORMATION: PS11: polymorphic base T or C  
1274 <220> FEATURE:  
1275 <221> NAME/KEY: misc\_feature  
1276 <222> LOCATION: (1261)..(1320)  
1277 <223> OTHER INFORMATION: n's represent sequence between PS11 and PS12  
1280 <220> FEATURE:  
1281 <221> NAME/KEY: allele  
1282 <222> LOCATION: (1350)..(1350)  
1283 <223> OTHER INFORMATION: PS12: polymorphic base C or T  
1286 <220> FEATURE:  
1287 <221> NAME/KEY: misc\_feature  
1288 <222> LOCATION: (1381)..(1440)  
1289 <223> OTHER INFORMATION: n's represent sequence between PS12 and PS13  
1292 <220> FEATURE:  
1293 <221> NAME/KEY: allele  
1294 <222> LOCATION: (1470)..(1470)  
1295 <223> OTHER INFORMATION: PS13: polymorphic base G or A  
1298 <220> FEATURE:  
1299 <221> NAME/KEY: misc\_feature  
1300 <222> LOCATION: (1501)..(1560)  
1301 <223> OTHER INFORMATION: n's represent sequence between PS13 and PS14  
1304 <220> FEATURE:  
1305 <221> NAME/KEY: allele  
1306 <222> LOCATION: (1590)..(1590)  
1307 <223> OTHER INFORMATION: PS14: polymorphic base C or T  
1310 <220> FEATURE:  
1311 <221> NAME/KEY: misc\_feature  
1312 <222> LOCATION: (1621)..(1680)  
1313 <223> OTHER INFORMATION: 's represent sequence between PS14 and PS15  
1316 <220> FEATURE:  
1317 <221> NAME/KEY: allele  
1318 <222> LOCATION: (1710)..(1710)  
1319 <223> OTHER INFORMATION: PS15: polymorphic base A or G  
1322 <220> FEATURE:  
1323 <221> NAME/KEY: misc\_feature  
1324 <222> LOCATION: (1741)..(1800)  
1325 <223> OTHER INFORMATION: n's represent sequence between PS15 and PS16

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1385	tcctggattt ggagtgggc aagctggta y gtgtcaatgt gcagcaaaat taatttagat	1020
W--> 1386	nnnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn	1080
1387	tatgtgtctg gagtgcttc aaaggaccay tgctcagagg aataacttgc cactacagg	1140
W--> 1388	nnnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn	1200
1389	tgaaaatgtc agagtttgt gcaacatady agtagcaga aaaaccaagt gaaaagtctt	1260
W--> 1390	nnnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn	1320
1391	gagaaggtac taaattgtt ggtatttccy gtaggaaccc cagagcgaaa tacagttgc	1380
W--> 1392	nnnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn	1440
1393	gcttgatttt attcaaactt tgcattttcatatttat cttggaaaat tcaattgtgt	1500
W--> 1394	nnnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn	1560
1395	gtagacaatt tgcctggcac caaagtaaay gcagagagt tagagaggat aaaacggcaa	1620
W--> 1396	nnnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn	1680
1397	aaagtaaacg cagagagtgt agagaggatr aaacggcaac acagctcaca agaacagact	1740
W--> 1398	nnnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn	1800
1399	caagaacaga ctttccagct gctgaagttr tggaaacatc aaaacaaaga ccaagatata	1860
W--> 1400	nnnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn	1920
1401	atagtcaga agatcatcca aggtatgatm atctaaaata aaaagatcaa tcagaaatca	1980
W--> 1402	nnnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn	2040
1403	atgcagatat tgacctctgt gaaaacagcr tgcagcggca cattggacat gctaaccctca	2100
W--> 1404	nnnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn	2160
1405	gcttcacaat gtacaaattt gatcagaagy tatttttaga aatgataggt aaccaggccc	2220
W--> 1406	nnnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn	2280
E--> 1409	Page 25	

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VERIFICATION SUMMARY  
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Input Set : A:\TNFRSF11B seq list.txt  
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L:13 M:270 C: Current Application Number differs, Replaced Current Application Number  
L:155 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1  
L:155 M:112 C: (48) String data converted to lower case,  
L:1370 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:94  
L:1372 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:94  
L:1374 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:94  
L:1376 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:94  
L:1378 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:94  
L:1380 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:94  
L:1382 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:94  
L:1384 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:94  
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L:1388 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:94  
L:1390 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:94  
L:1392 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:94  
L:1394 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:94  
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L:1398 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:94  
L:1400 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:94  
L:1402 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:94  
L:1404 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:94  
L:1406 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:94  
L:1409 M:254 E: No. of Bases conflict, LENGTH:Input:25 Counted:2284 SEQ:94  
L:1409 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:2  
L:1409 M:252 E: No. of Seq. differs, <211>LENGTH:Input:2280 Found:2284 SEQ:94